

IFWO

RAW SEQUENCE LISTING

DATE: 07/21/2004 TIME: 09:40:36

PATENT APPLICATION: US/10/786,380

Input Set : A:\2316.2009-000SEQLIST.TXT
Output Set: N:\CRF4\07212004\J786380.raw

```
4 <110> APPLICANT: Cardosa, Mary Jane
 5
        Sutter, Gerd
        Erfle, Volker
 8 <120> TITLE OF INVENTION: Recombinant MVA Virus Expressing Dengue
        Virus Antigens and the Use Thereof in Vaccines
12 <130> FILE REFERENCE: 2316.2009-000
14 <140> CURRENT APPLICATION NUMBER: US 10/786,380
15 <141> CURRENT FILING DATE: 2004-02-24
                                                              ENTERED
17 <150> PRIOR APPLICATION NUMBER: US 09/147,919
18 <151> PRIOR FILING DATE: 1999-03-23
20 <160> NUMBER OF SEQ ID NOS: 8
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 33
26 <212> TYPE: DNA
27 <213> ORGANISM: Unknown
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Primer for left 600 bp DNA flanking the site of a
         2500 bp deletion in the Hind III N fragment of MVA
31
         genome
34 <400> SEQUENCE: 1
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37 <210> SEQ ID NO: 2
38 <211> LENGTH: 42
39 <212> TYPE: DNA
40 <213> ORGANISM: Unknown
42 <220> FEATURE:
43 <223> OTHER INFORMATION: Primer for left 600 bp DNA flanking the site of a
         2500 bp deletion in the Hind III N fragment of MVA
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         genome
47 <400> SEQUENCE: 2
48 cagcageceg ggtattegat gattattttt aacaaaataa ca
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50 <210> SEQ ID NO: 3
51 <211> LENGTH: 36
52 <212> TYPE: DNA
53 <213> ORGANISM: Unknown
55 <220> FEATURE:
56 <223> OTHER INFORMATION: Primer for right 550 bp DNA flanking the site of a
57
         2500 bp deletion in the Hind III N fragment of MVA
58
        genome
60 <400> SEQUENCE: 3
61 cagcagctgc aggaatcatc cattccactg aatagc
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63 <210> SEQ ID NO: 4

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- 64 <211> LENGTH: 36
- 65 <212> TYPE: DNA

12.

- 66 <213> ORGANISM: Unknown
- 68 <220> FEATURE:
- 69 <223> OTHER INFORMATION: Primer for right 550 bp DNA flanking the site of a
- 2500 bp deletion in the Hind III N fragment of MVA
- 71 genome
- 73 <400> SEQUENCE: 4
- 74 caqcaqqcat qccqacqaac aaqqaactqt aqcaqa

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- 76 <210> SEQ ID NO: 5
- 77 <211> LENGTH: 33
- 78 <212> TYPE: DNA
- 79 <213> ORGANISM: Unknown
- 81 <220> FEATURE:
- 82 <223> OTHER INFORMATION: Primer for amplification of E. coli LacZ orf
- 84 <400> SEQUENCE: 5
- 85 cagcaggtcg accccgaccg ccttactgcc gcc

- 87 <210> SEQ ID NO: 6
- 88 <211> LENGTH: 33
- 89 <212> TYPE: DNA
- 90 <213> ORGANISM: Unknown
- 92 <220> FEATURE:
- 93 <223> OTHER INFORMATION: Primer for amplification of E. coli LacZ orf
- 95 <400> SEQUENCE: 6
- 96 ggggggctgc agatggtagc gaccggcgct cag

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- 98 <210> SEQ ID NO: 7
- 99 <211> LENGTH: 36
- 100 <212> TYPE: DNA
- 101 <213> ORGANISM: Unknown
- 103 <220> FEATURE:
- 104 <223> OTHER INFORMATION: Primer for isolation of cDNA fragment of Dengue
- virus 2NGC strain 105
- 107 <400> SEQUENCE: 7
- 108 cagcagcccg ggatggcagc aatcctggca tacacc

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- 110 <210> SEQ ID NO: 8
- 111 <211> LENGTH: 39
- 112 <212> TYPE: DNA
- 113 <213> ORGANISM: Unknown
- 115 <220> FEATURE:
- 116 <223> OTHER INFORMATION: Primer for isolation of cDNA fragment of Dengue
- virus 2NGC strain 117
- 119 <400> SEQUENCE: 8
- 120 cagcageceg ggtcactgta gtttgtccat ceteagect

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VERIFICATION SUMMARY

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